

fanjin

July 8, 2021

lets follow Fanjin a little more closely /home/dan/usb_twitching/library/core/pnas_Wong_2011_slingshot_mo
the purpose of this is to get a better understanding of analysing the trajectory so that we can fully
understand how our simulated data differs from the experimental data

```
[1]: import os
import sys
import collections
import numpy as np
import _fj
import matplotlib.pyplot as plt
import matplotlib as mpl
import command
import readtrack
import twanalyse
from tqdm import tqdm
```

```
[2]: verbose = False
notename = 'fanjin'
notedir = 'fanjin/'
if not os.path.exists(notedir):
    os.makedirs(notedir)
join = os.path.join
def _savefig(name):
    d = os.path.dirname(name)
    if d:
        if not os.path.exists(d):
            os.makedirs(d)
    save = join(notedir, name)
    print("saving to ", save)
    plt.savefig(save)
```

```
[3]: # -----
idx, ltrs = _fj.slicehelper.load_linearized_trs('all')
```

100%| | 3113/3113 [00:01<00:00, 2709.50it/s]

```
[4]: crawling_idx = _fj.slicehelper.load('default_crawling_list')
crawling = [ltrs[i] for i in crawling_idx]
```

```

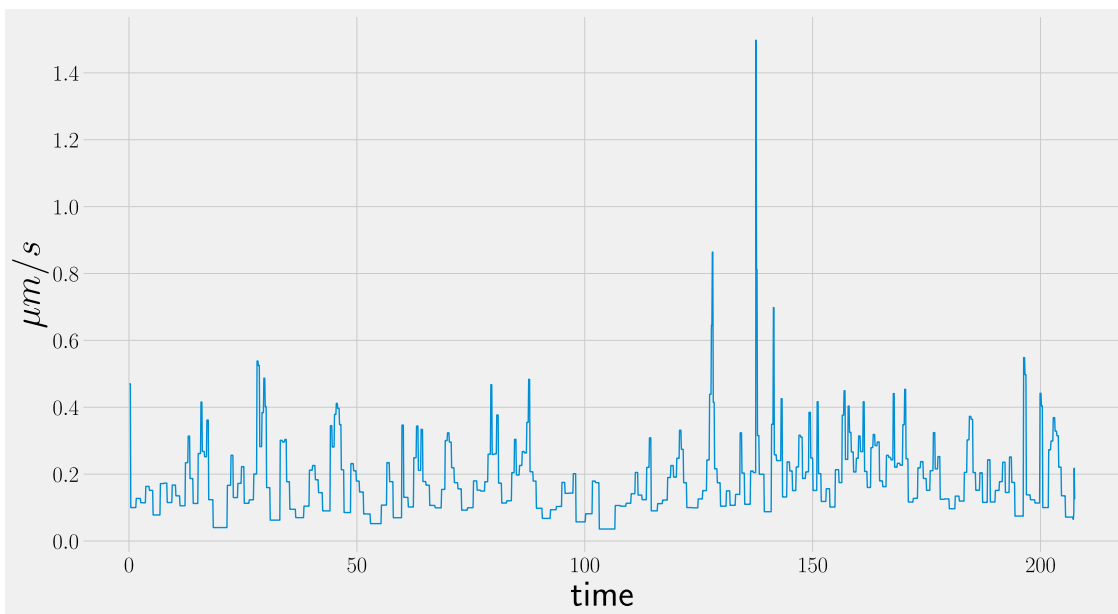
candidate_idx = 2924
candidate = ltrs[candidate_idx]
whitelist = _fj.slicehelper.load('candidates_whitelist')
whitelist_tr = [ltrs[i] for i in whitelist]
# -----

```

```

[5]: # -----
ax = plt.gca()
def plot_candidate(ax, track):
    time = track['time']
    velocity = track.get_head_v()
    track_speed = np.linalg.norm(velocity, axis=1)
    ax.plot(time[1:]-time[0], track_speed)
    ax.set_xlabel('time')
    ax.set_ylabel('$\mu\text{m/s}$')
# plot_candidate(ax, ltrs[10])
plot_candidate(ax, candidate)
plt.show()

```



these are linearised tracks, lets check that displacements add up to $0.12\ \mu\text{m}$

```

[6]: # plot displacement angles, body axis angles
# for individual linearised trajectories

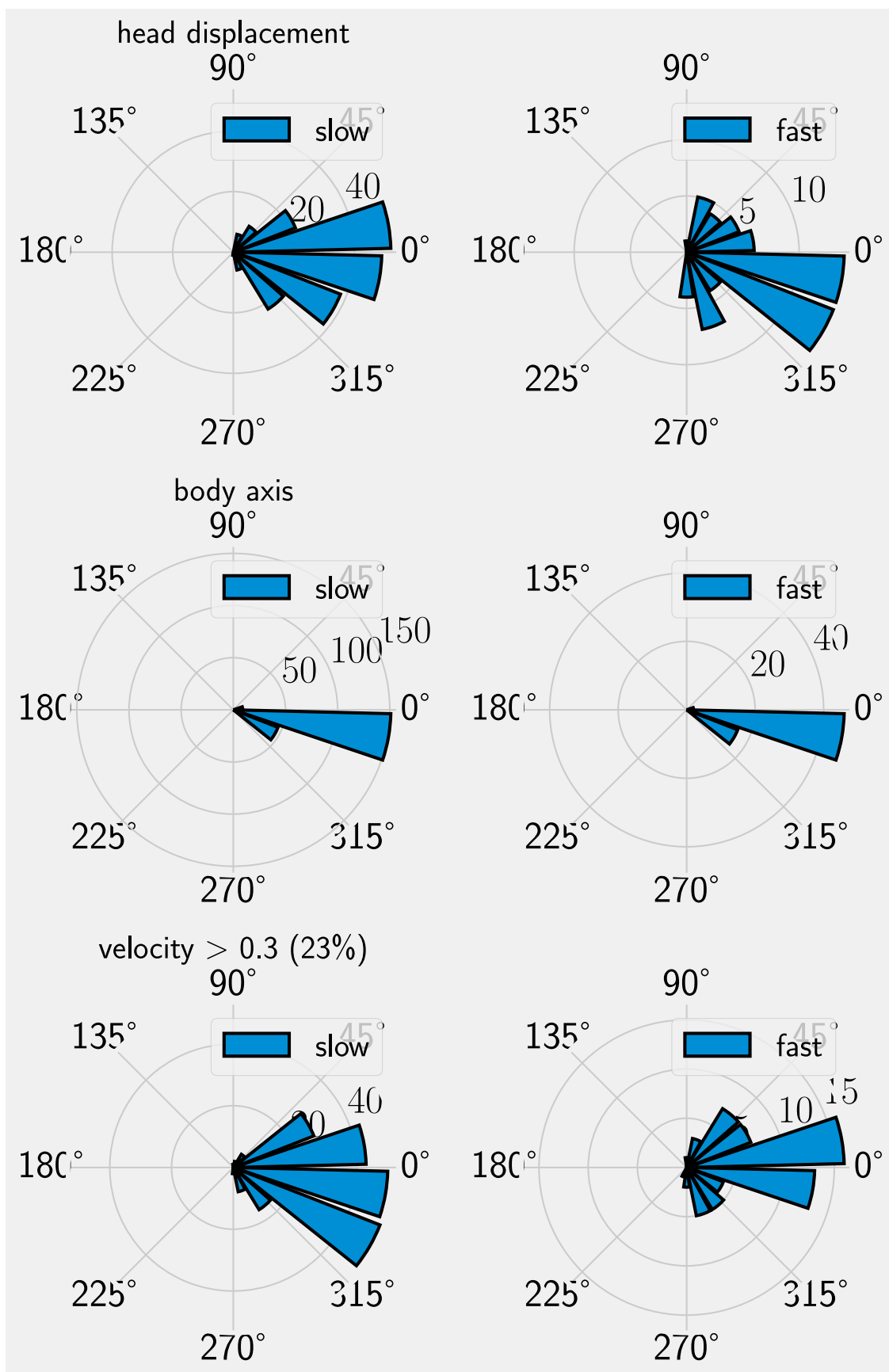
allpolar = twanalyse.allpolar
plotpolar = twanalyse.plotpolar
def plotpolar_with_axes(polardata):

```

```
fig, axes = plt.subplots(3, 2, subplot_kw=dict(polar=True),
                        figsize=(8,12))
plotpolar(axes, polardata)
```

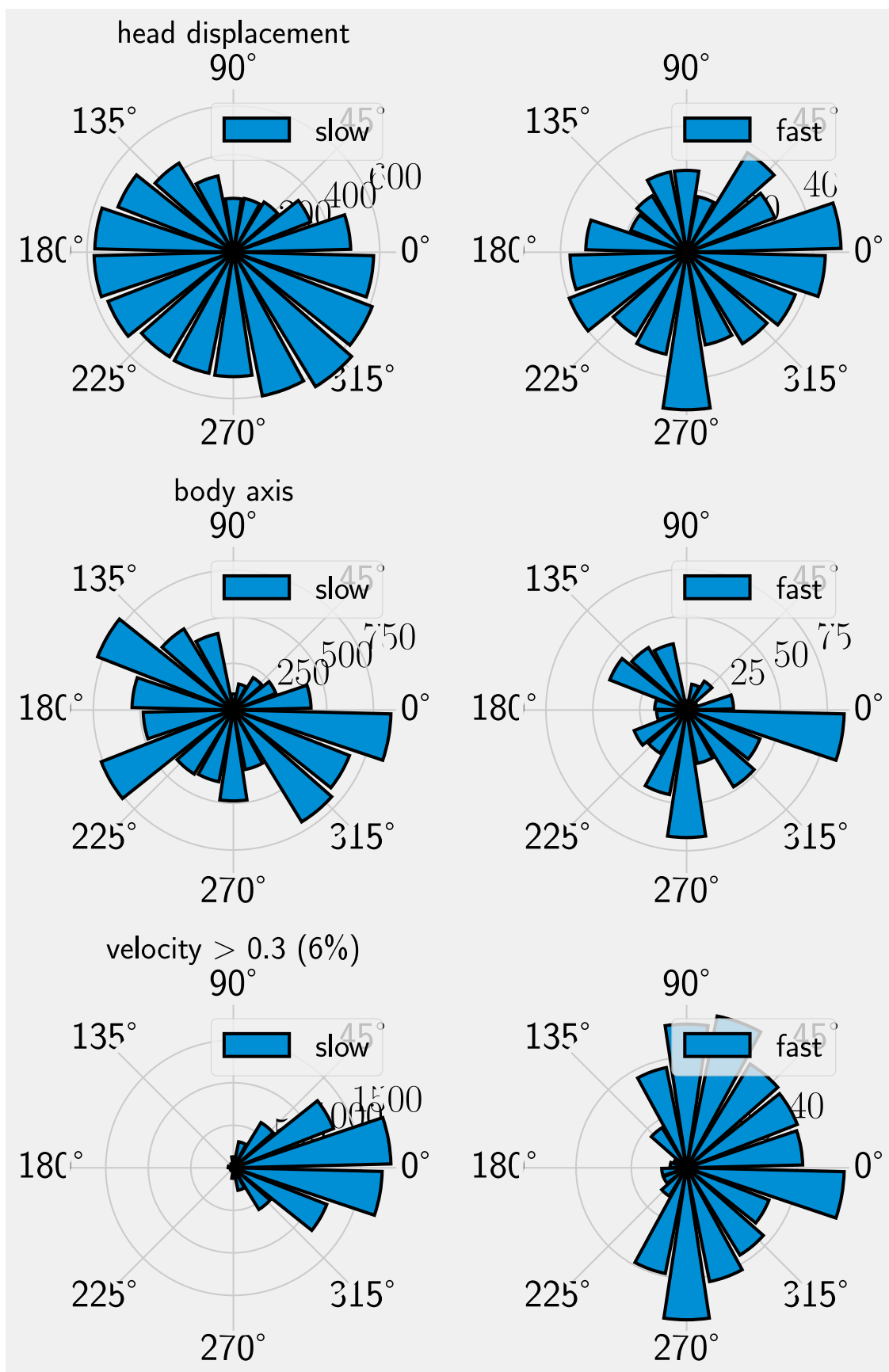
```
[7]: # polar plots for the candidate track only
polardata = allpolar([candidate])
plotpolar_with_axes(polardata)
_savefig("polar_candidate.png")
```

```
100%|      | 1/1 [00:00<00:00, 1305.42it/s]
saving to  fanjin/polar_candidate.png
```



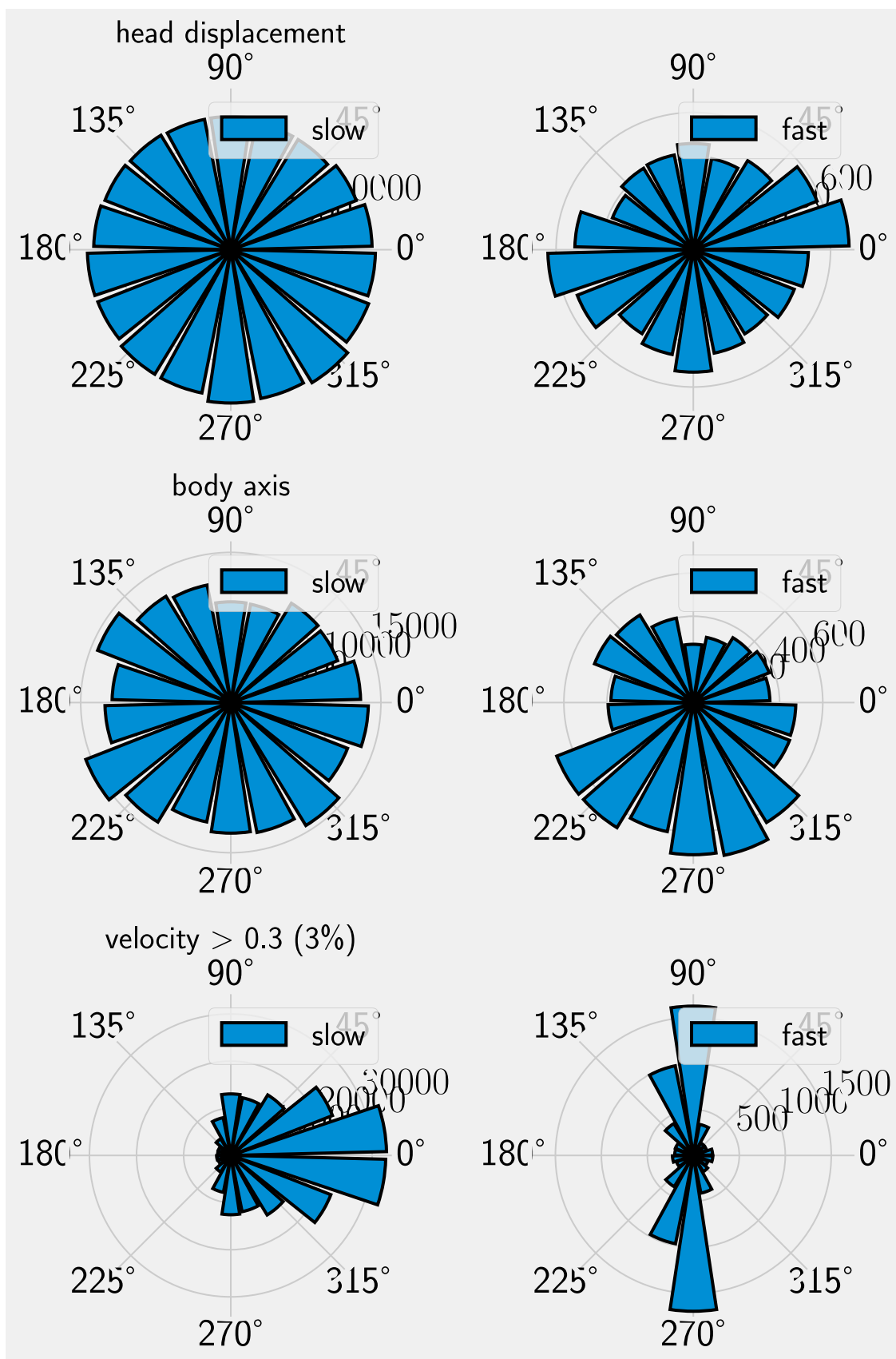
```
[8]: # polar plots for whitelist
polardata = allpolar(whitelist_tr)
plotpolar_with_axes(polardata)
_savefig("polar_whitelist.png")
```

```
100%|      | 64/64 [00:00<00:00, 3980.30it/s]
saving to  fanjin/polar_whitelist.png
```



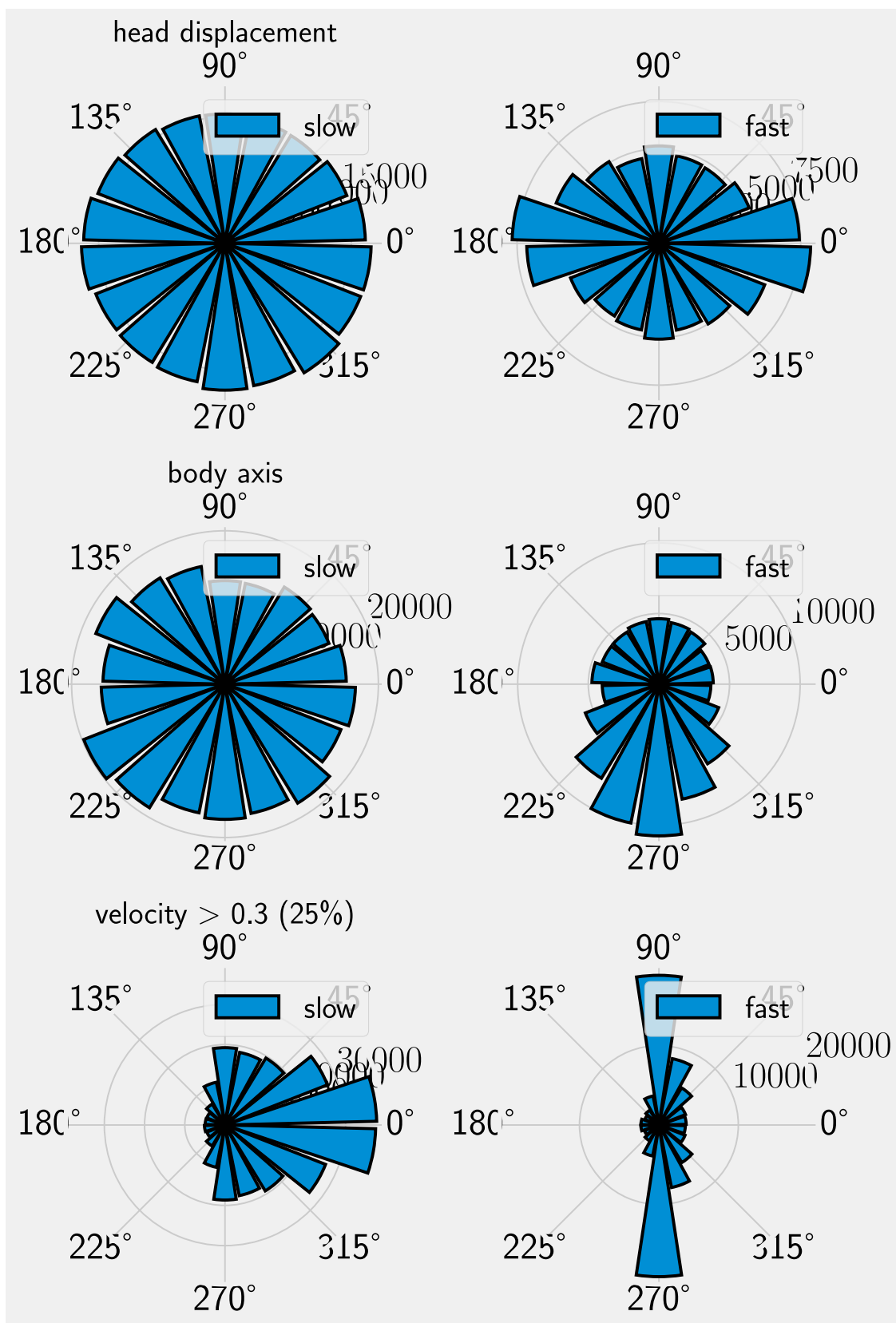
```
[9]: # crawling data
polardata = allpolar(crawling)
plotpolar_with_axes(polardata)
_savefig("polar_crawling.png")
```

```
100%|      | 2505/2505 [00:00<00:00, 2880.64it/s]
saving to  fanjin/polar_crawling.png
```




```
[10]: # all data
polardata = allpolar(ltrs)
plotpolar_with_axes(polardata)
_savefig("polar_all.png")

0%|          | 0/3113 [00:00<?,
?it/s]/home/dan/usb_twitching/pili/src/analysis/twanalyse.py:803:
RuntimeWarning: invalid value encountered in true_divide
    norm_dy = dy/np.linalg.norm(dy, axis=1)[: ,np.newaxis]
100%|         | 3113/3113 [00:01<00:00, 2671.21it/s]
saving to  fanjin/polar_all.png
```



```

[11]: # velocity > 0.3 (25%) ???
# is this a bug in my code or due outliers in the 'all' data
# lets make a walking list with a mean velocity threshold (?)
# see localdir for the removed outliers
localdir = 'vel_outlier'

def velocity_outliers(idx, ltrs, minvel=0.0, maxvel=1.0):
    meanvel = np.array([np.mean(np.linalg.norm(ltr.
    ↳get_step_velocity(),axis=1)) for ltr in ltrs])
    print(np.isnan(meanvel).shape)
    print(idx.shape)
    nan_idx = idx[np.isnan(meanvel)]
    left_idx = idx[meanvel < minvel]
    right_idx = idx[meanvel > maxvel]
    return left_idx, right_idx, nan_idx

left, right, nan = velocity_outliers(idx, ltrs, maxvel=1.0)
bad_idx = np.union1d(right, nan)
l, r, n = left.size, right.size, nan.size
print('left\t{:4d}\tright\t{:4d}\tnan\t{:4d}'.format(l, r, n))
print('so there are {:d} candidates to remove'.format(bad_idx.size))

if verbose:
    import shapeplot
    for track_id in right:
        ltrack = ltrs[track_id]
        speed = np.linalg.norm(ltrack.get_step_velocity(), axis=1)
        print('fast step proportion', npcount(speed > 0.3)/speed.size)
        ax = plt.gca()
        shapeplot.longtracks(ax,[ltrack])
        name = join(localdir,'lt_{:04d}.png')
        _savefig(name.format(track_id))
        ax.clear()
    plt.close()

safeall = np.delete(idx, bad_idx)

```

```

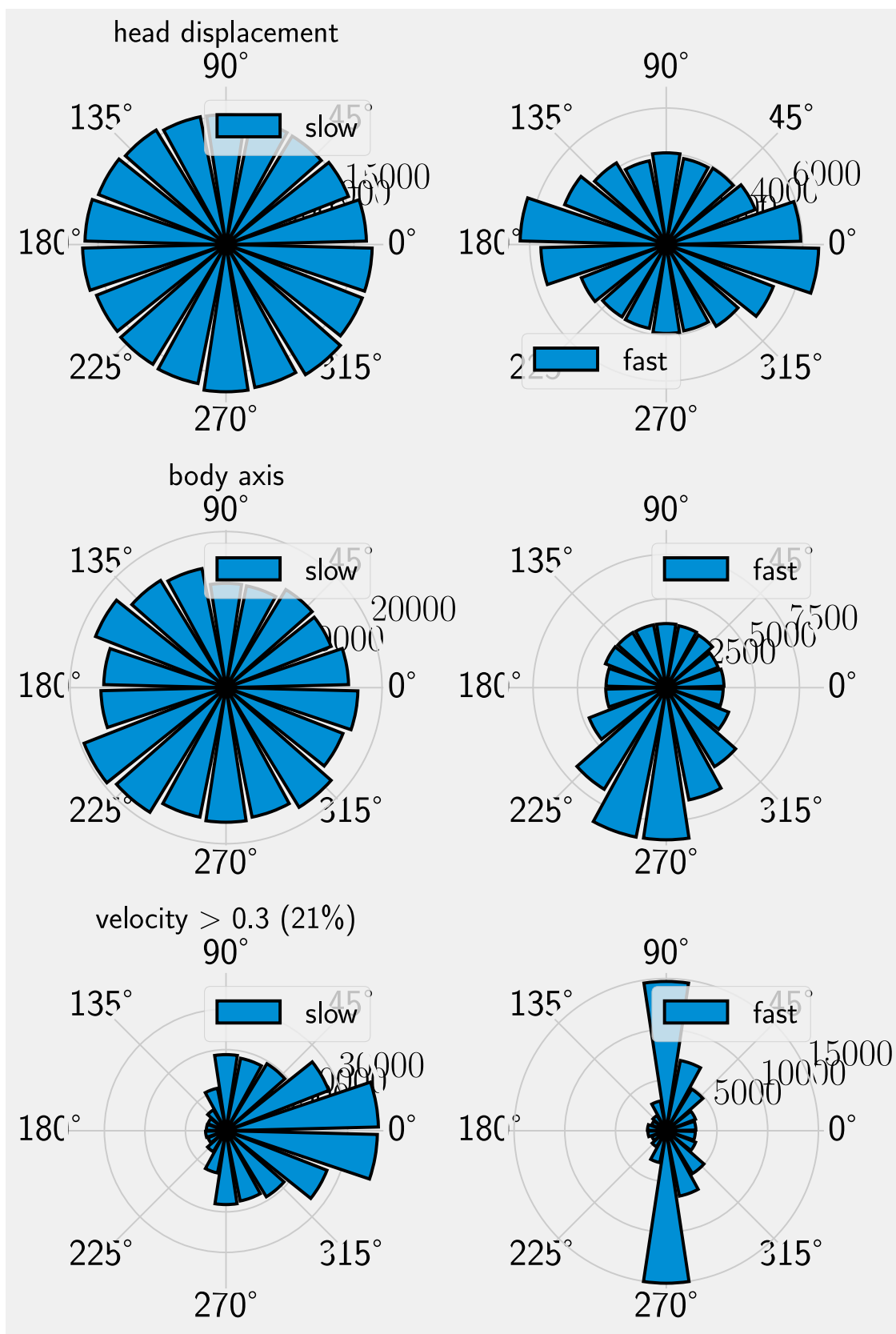
(3113,)
(3113,)
left      0    right 30      nan   6
so there are 36 candidates to remove
/home/dan/.local/lib/python3.8/site-packages/numpy/core/fromnumeric.py:3419:
RuntimeWarning: Mean of empty slice.
    return _methods._mean(a, axis=axis, dtype=dtype,
/home/dan/.local/lib/python3.8/site-packages/numpy/core/_methods.py:188:
RuntimeWarning: invalid value encountered in double_scalars

```

```
ret = ret.dtype.type(ret / rcount)
```

```
[12]: polardata = allpolar([ltrs[i] for i in safeall])
      plotpolar_with_axes(polardata)
      _savefig("polar_all.png")
      # and the value did change from 25% to 21% but its still huge
      # compared to the crawling set, need more work to see if
      # high speed individuals in walking set are well resolved or not.
```

```
100%|      | 3077/3077 [00:01<00:00, 2943.91it/s]
saving to  fanjin/polar_all.png
```

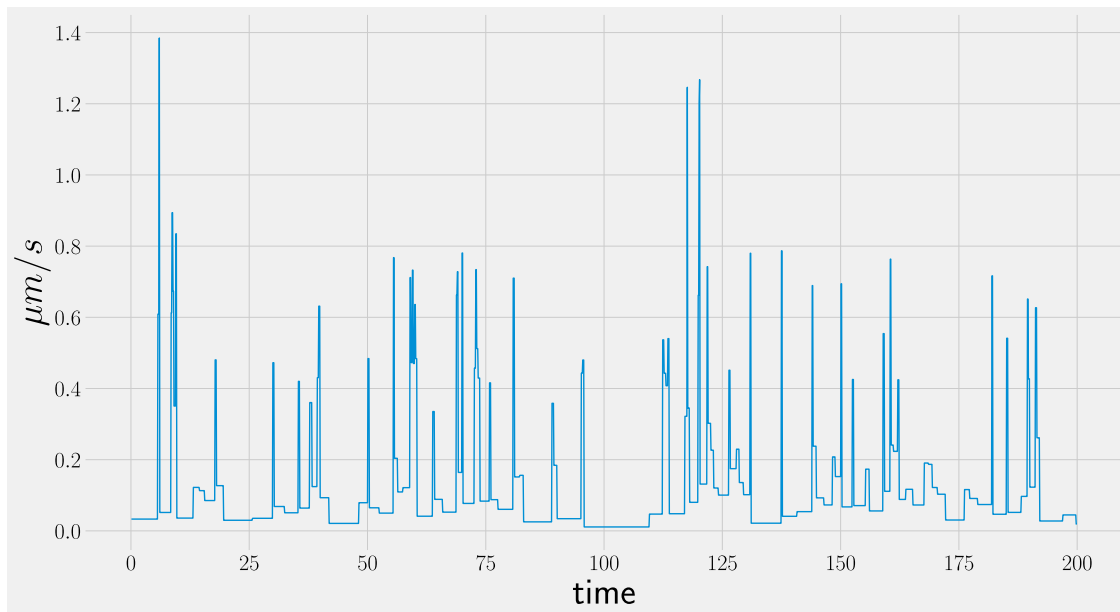


```
[13]: # * candidate trajectory, all polar plots are oriented towards direction of
      ↪ motion
      # * whitelist, slow actions are polarised, fast actions have large components
      ↪ in forward direction
      # and also at 90 degrees
      # * all tracks, the deviation angle for the fast mode are now dominated by
      # 90 degree rotations, so these are associated with slower moving trajectories
```

```
[14]: # -----
      # lets do the same on simulation dataset
      simdir = "/home/dan/usb_twitching/run/new/angle_smoothed/range_pbrf/
      ↪ anchor_angle_smoothing_fraction_00.250"
      with command.chdir(simdir):
          simtrs = readtrack.trackset()
      sim_ltrs = [_fj.linearize(tr) for tr in simtrs]
      simtr = sim_ltrs[0]

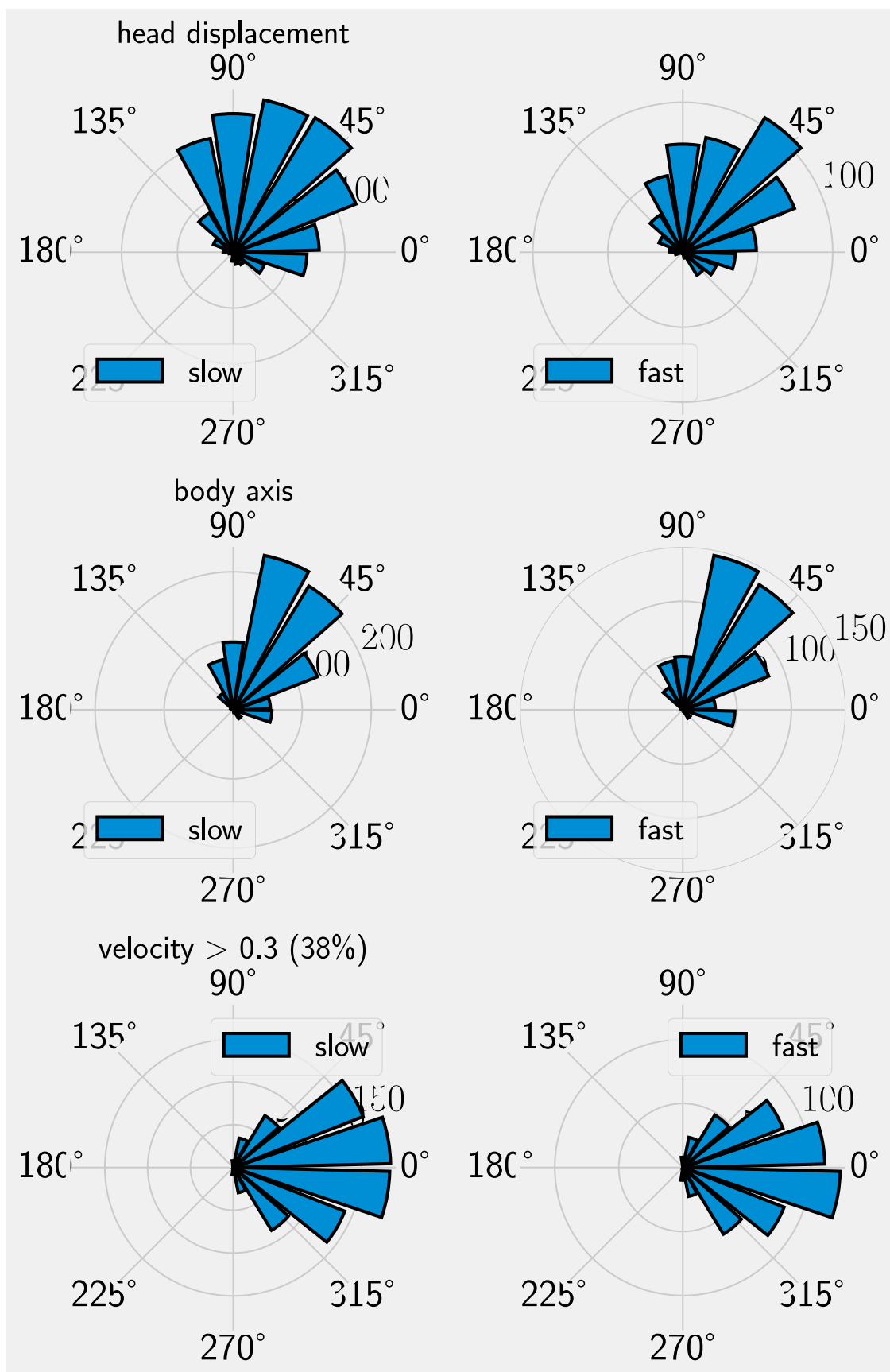
      plot_candidate(plt.gca(), simtr.part(0,2000))
```

searching for tracks with form data/bacterium_*.dat



```
[15]: # simulated data
      polardata = allpolar(sim_ltrs)
      plotpolar_with_axes(polardata)
```

100% | 1/1 [00:00<00:00, 589.34it/s]

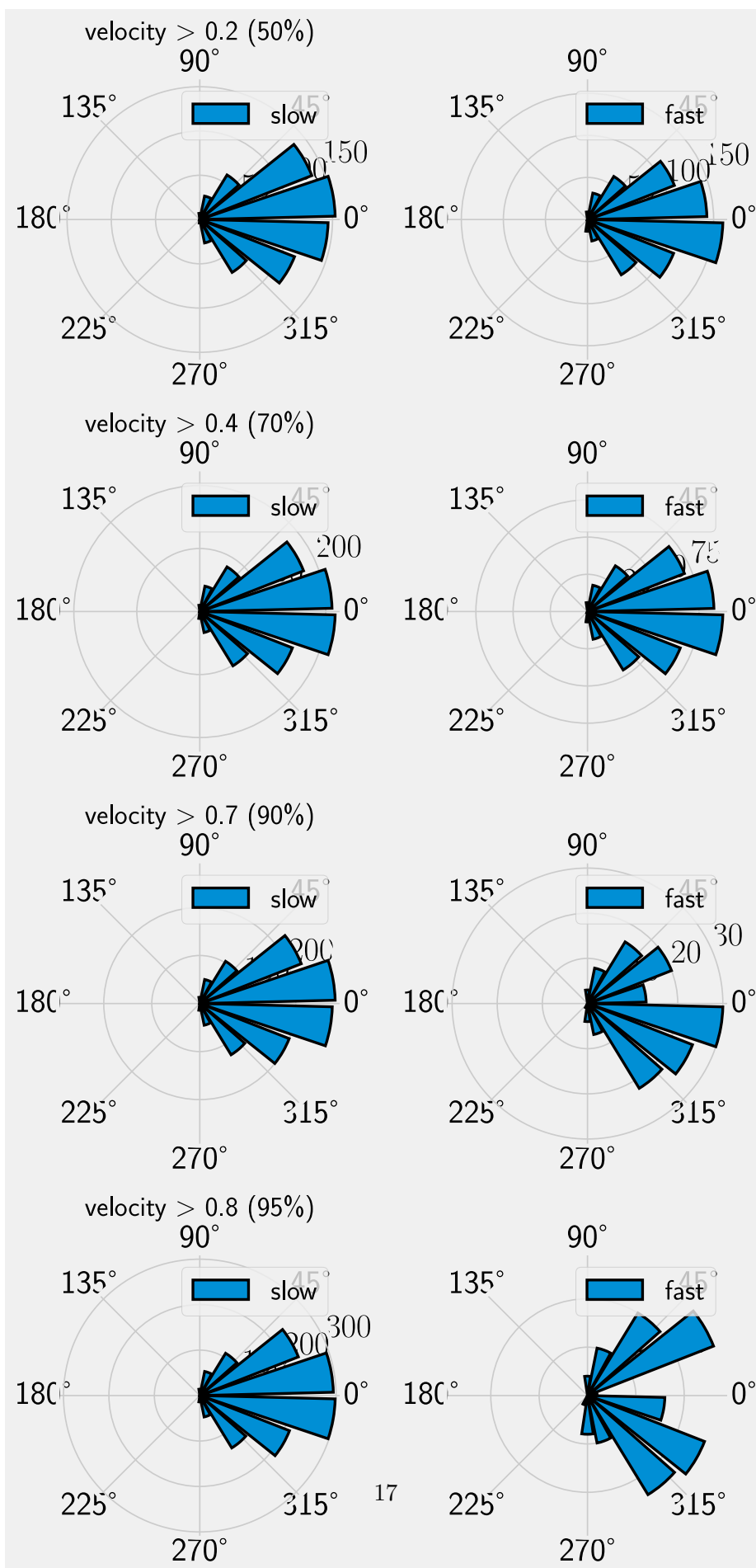


which is similar to candidate. but note that there is not much to distinguish fast and slow.

```
[16]: # what about using a different velocity threshold?
# can we see a difference between fast and slow then?
vthreshold = 0.3
quantiles = [0.5, 0.7, 0.9, 0.95]
actdata = twanalyse.actions(sim_ltrs)
def fraction(x, threshold):
    npcount = np.count_nonzero
    return npcount(x>threshold)/x.size

N = len(quantiles)
fig, axes = plt.subplots(N, 2, subplot_kw=dict(polar=True),
    figsize=(8,4*N))
for i, q in enumerate(quantiles):
    quant = np.quantile(actdata['velocity'], q)
    row = axes[i]
    pdata = twanalyse.allpolar(sim_ltrs, quant)
    fastidx = actdata['velocity'] > quant
    twanalyse.polar_dev(row, pdata.deviation, fastidx)
    row[0].set_title(r'veLOCITY $>$ {:.3.1f} ({:d}\%)'.format(
        quant, int(100*q)
    ))
plt.tight_layout()
```

```
100%|      | 1/1 [00:00<00:00, 1565.04it/s]
100%|      | 1/1 [00:00<00:00, 601.51it/s]
100%|      | 1/1 [00:00<00:00, 583.11it/s]
100%|      | 1/1 [00:00<00:00, 587.52it/s]
100%|      | 1/1 [00:00<00:00, 579.32it/s]
```

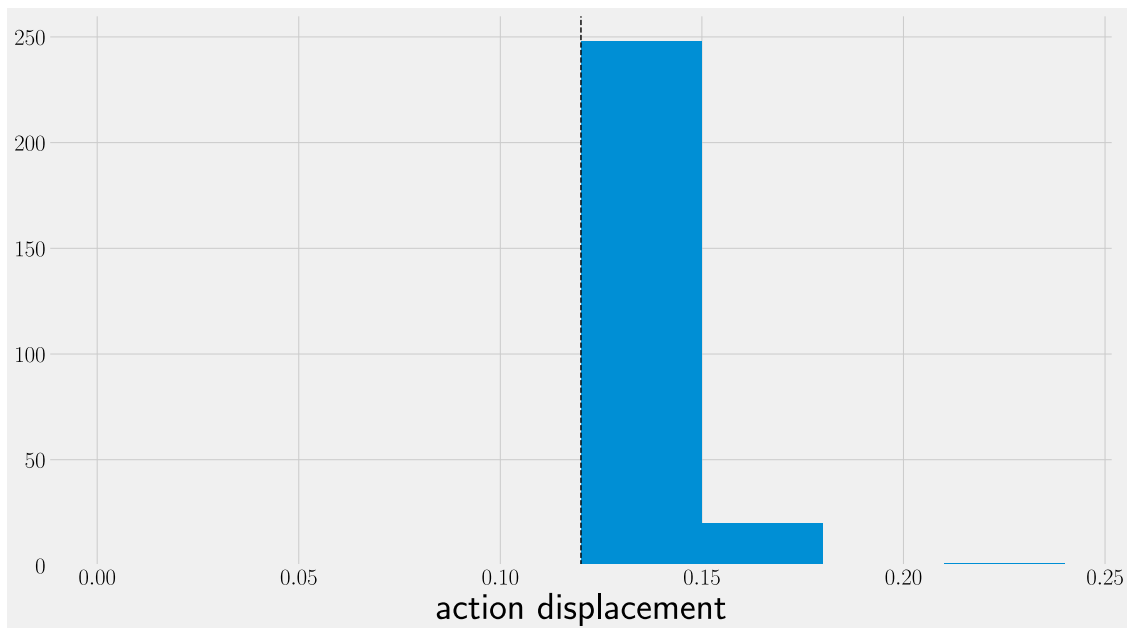



```
[17]: candidate_action = twanalyse.actions([candidate])
```

100%| | 1/1 [00:00<00:00, 3435.14it/s]

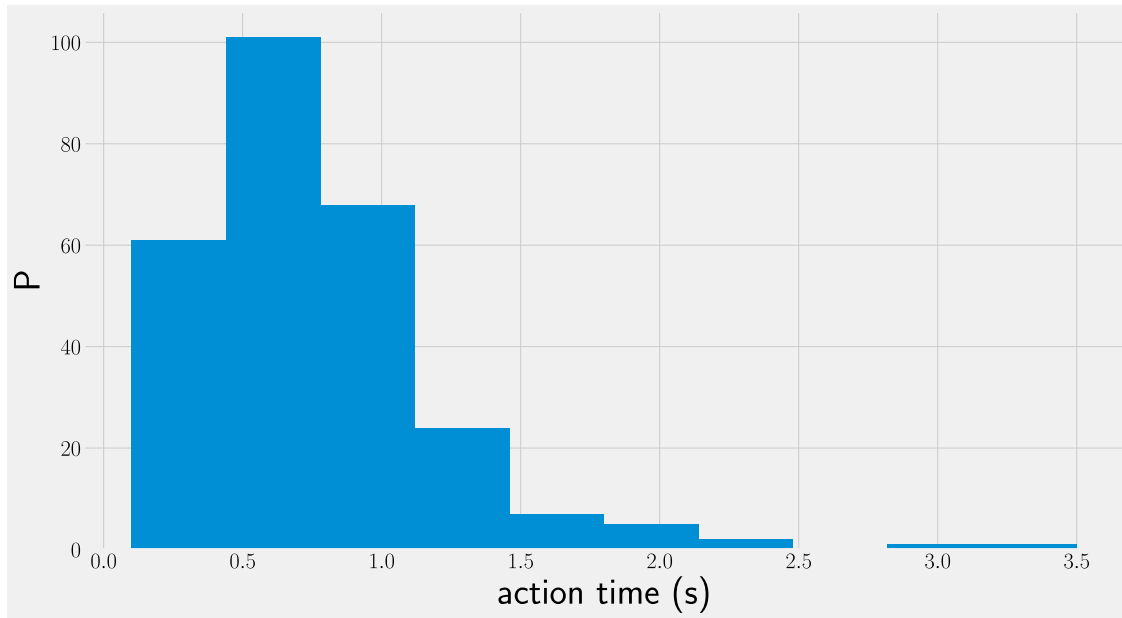
```
[18]: # plot candidate displacements
ax = plt.gca()
bins = np.arange(0,max(candidate_action['dx']),0.03)
ax.hist(candidate_action['dx'], bins=bins)
ax.set_xlabel('action displacement')
ax.axvline(0.12, linestyle='--', color='k')
```

```
[18]: <matplotlib.lines.Line2D at 0x7f3934dde760>
```



```
[19]: # plot candidate action dt
ax = plt.gca()
ax.set_ylabel('P')
ax.set_xlabel('action time (s)')
ax.hist(candidate_action['dt'])
```

```
[19]: (array([ 61., 101., 68., 24., 7., 5., 2., 0., 1., 1.]),
array([0.1 , 0.44, 0.78, 1.12, 1.46, 1.8 , 2.14, 2.48, 2.82, 3.16, 3.5 ]),
<BarContainer object of 10 artists>)
```

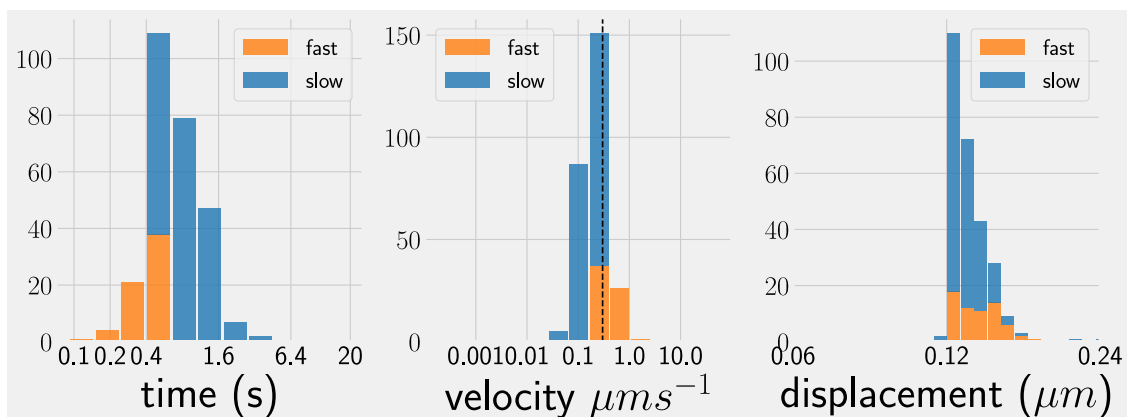


```
[20]: actions = twanalyse.actions
      plot_actiondata = twanalyse.plot_actiondata
      candidate_action = actions([candidate])
      traction = actions(crawling)
```

```
100%|      | 1/1 [00:00<00:00, 3086.32it/s]
100%|      | 2505/2505 [00:00<00:00, 22919.99it/s]
```

```
[21]: plot_actiondata(candidate_action)
      _savefig('candidate_actions.png')
      plt.show()
```

```
trimming 0.0 elements < 0 && 0.0 elements > 100
trimming 0.0 elements < 0 && 0.0 elements > 100
saving to fanjin/candidate_actions.png
```



```
[22]: # all data
if verbose:
    plot_actiondata(traction)
    _savefig('actions.png')
    plt.show()

[23]: # [markdown]
# find out where the 0.1/0.2 second fast actions are hiding, look at those
→ trajectories by eye.

[24]: # plot 2d histogram for actions A,A+1

def action_corr(actions):
    # first bin all the actions but keep track of which bin they belong to
    vel = actions['velocity']
    vel = twanalyse.trim_limits(vel, [0.004,16])
    bins = np.geomspace(0.004,33,16+1,True)
    with np.printoptions(precision=3, suppress=True):
        print('bins', bins)
    bincount = np.zeros(bins.size-1,dtype=int) # one less bin than edge
    binmap = []
    nbin = []
    count2d = np.zeros((bins.size-1,bins.size-1), dtype=int)
    for i in range(vel.size-1):
        v = vel[i]
        v_n = vel[i+1]
        bin_idx = np.searchsorted(bins, v)
        nbin_idx = np.searchsorted(bins, v_n)
        binmap.append(bin_idx)
        nbin.append(nbin_idx)
        bincount[bin_idx] += 1
        count2d[bin_idx][nbin_idx] += 1

    X = bins
    Y = bins
    return X, Y, count2d

X, Y, count2d = action_corr(candidate_action)
def plot_c2d(X, Y, count2d):
    ax = plt.gca()
    ax.set_xscale('log')
    ax.set_yscale('log')
    cmap = mpl.cm.get_cmap()
    cmlist = [cmap(c) for c in np.linspace(0,1,100,True)]
    cm = mpl.colors.LinearSegmentedColormap.from_list('adj', cmlist)
```

```

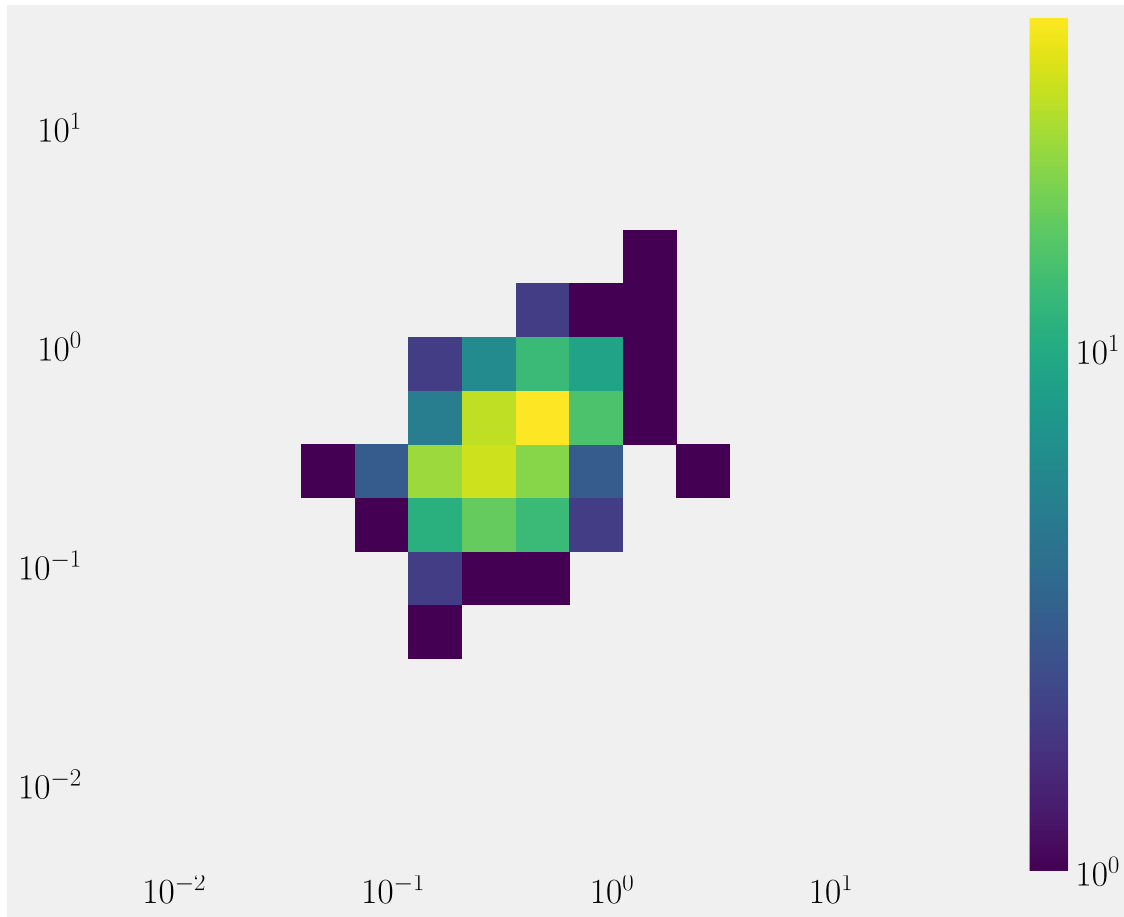
Im = np.ma.masked_where(count2d == 0, count2d)
# norm = mpl.colors.Normalize(count2d.min(), count2d.max())
norm = mpl.colors.LogNorm(1.0, count2d.max())
pos = plt.pcolormesh(X, Y, Im, cmap=cm, norm=norm)
plt.colorbar(pos)
ax.set_aspect('equal')
plot_c2d(X, Y, count2d)

```

```

trimming 0.0 elements < 0.004 && 0.0 elements > 16
bins [ 0.004  0.007  0.012  0.022  0.038  0.067  0.118  0.207  0.363  0.638
       1.122  1.971  3.463  6.084 10.689 18.782 33.   ]

```



```

[25]: X, Y, c2d = action_corr(traction)
      plot_c2d(X, Y, c2d)

```

```

trimming 0.04453258316860121 elements < 0.004 && 4.223900518694984e-06 elements
> 16
bins [ 0.004  0.007  0.012  0.022  0.038  0.067  0.118  0.207  0.363  0.638
       1.122  1.971  3.463  6.084 10.689 18.782 33.   ]

```

